



FIG S3 Inactivation of *acrT* in *Sac. erythraea*. (A) Schematic diagram of *acrT* deletion by homologous recombination with the linearized fragment in *Sac. erythraea* A226. (B) Identification of *acrT* deletion in *Sac. erythraea* A226. M, 5,000 bp DNA ladder; pUCTSR Δ *acrT*, the positive control, from which 1,870 bp DNA fragment was amplified; A226, the negative control, from which 756 bp DNA fragment was amplified; A226 Δ *acrT*, the screened mutant, from which 1,870 bp DNA fragment was amplified. (C) Aerial mycelia formation of A226 and its derivatives. All strains were grown on R3M solid medium at

8 30 °C for 48 and 72 h. (D) Growth curves of A226 and A226 Δ *acrT* in R5 liquid medium. Their dry
9 weights of mycelia (DWM) were measured. (E) Confirmation of *acrT* deletion in the industrial strain *Sac.*
10 *erythraea* WB. M, 5,000 bp DNA ladder; pUCTSR Δ *acrT*, the positive control, from which 1,870 bp DNA
11 fragment was amplified; WB, the negative control, from which 756 bp DNA fragment was amplified;
12 WB Δ *acrT*, the screened mutant, from which 1,870 bp DNA fragment was amplified. (F) Er-A production
13 in WB and WB Δ *acrT*. Mean values of $n = 3$ measurements are shown with SDs. *, $P < 0.05$.